CLAIM AMENDMENTS

(Insertions indicated by underline; deletions indicated by strikethrough.)

- 1. (Currently Amended) A recombinant expression vector comprising consisting essentially of an open reading frame operably linked to one or more regulatory elements, wherein the open reading frame encodes a of a nucleotide sequence encoding the polypeptide set forth in SEQ ID NO: 5.
- 2. (Currently Amended) The recombinant expression vector of Claim 1, wherein said open reading frame has nucleotide sequence is the nucleotide sequence set forth in SEQ ID NO: 4.
- 3. (Currently Amended) The recombinant expression vector of Claim 1, wherein said vector further comprises a replication-defective virus.
- 4. (Original) A host cell comprising the recombinant expression vector of Claim 1, wherein said host cell is selected from the group consisting of prokaryotic host cells and eukaryotic host cells.
 - 5. Cancelled.
- 6. (Currently Amended) A method for detecting <u>a</u> nucleic acids encoding Rig in a sample, comprising the steps of:
 - a) providing:
 - i) a sample comprising a nucleic acid encoding Rig,
 - ii) a <u>nucleic acid</u> probe comprising nucleic acid having complementarity to at least a portion of the nucleotide sequence of SEQ ID NO:4,
 - b) combining said sample and said probe under conditions wherein a hybridization complex is formed between said probe and said nucleic acid in said sample, and
- c) detecting said hybridization complex, whereupon the detection of the hybridization complex indicates the presence of a nucleic acid encoding Rig in the sample.

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- 7. (Original) The method of Claim 6, wherein said sample is selected from the group consisting of total cellular RNA, polyA RNA and genomic DNA.
 - 8. (Currently Amended) The method of Claim 6, wherein said sample comprises is from tumor tissue.
 - 9. (Original) The method of Claim 6, wherein said sample is from a human subject.
- 10. (Currently Amended) The method of Claim 6, wherein said method eomprises hybridization complex in step c) is detected using a Northern blotting protocol.
- 11. (Currently Amended) A method for amplifying <u>a</u> nucleic acids encoding Rig in a sample, comprising:
 - a) providing:
 - i) a sample comprising a nucleic acids encoding Rig,
 - ii) a DNA polymerase;
 - iii) two oligonucleotides, one of which is complementary having complementarity to the nucleotide sequence of SEQ ID NO:4 and one of which is complementary to the nucleotide sequence that is complementary to SEQ ID NO: 4; and
 - iv) <u>polymerase chain reaction (PCR)</u> amplification reagents;
 - b) combining said sample, said DNA polymerase, said oligonucleotides, and said PCR amplification reagents;
 - c) annealing said oligonucleotides to said nucleic acid in said sample; and
 - extending said oligonucleotides with reiterated DNA synthesis under conditions such that said nucleic acid is amplified, whereupon a nucleic acid encoding Rig is amplified. to produce an amplified product; and
 - e) detecting said amplified product.

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- 12. (Original) The method of Claim 11, wherein said DNA polymerase has both DNA-dependent DNA polymerase activity and reverse transcriptase RNA-dependent DNA polymerase activity.
 - 13. (Original) The method of Claim 11, wherein said sample is from a human subject.
 - 14. (Currently Amended) The method of Claim 11, wherein said sample comprises is from tumor tissue.
- 15. (Original) The method of Claim 11, wherein said nucleic acid is selected from DNA and RNA.
- 16. (Currently Amended) The method of Claim 11, wherein <u>one of said two</u> oligonucleotides <u>consists of emprise SEQ ID NO:2</u> and <u>the other of said two</u> <u>oligonucleotides consists of SEQ ID NO:3</u>.

17-28. (Cancelled)

29. (New) The method of claim 11, wherein the method further comprises step e) detecting said amplified product.

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